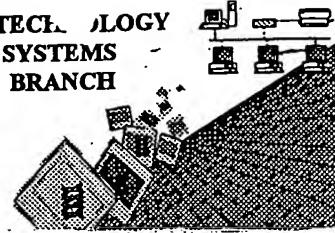


RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/027,736
Source: 01PE
Date Processed by STIC: 5/13/02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Raw Sequence Listing Error Summary

O1PE

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u> <u>101027,736</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <input type="checkbox"/> Wrapped Nucleic Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <input type="checkbox"/> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <input type="checkbox"/> Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 <input type="checkbox"/> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <input type="checkbox"/> Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <input type="checkbox"/> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <input type="checkbox"/> Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <input type="checkbox"/> Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If Intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <input checked="" type="checkbox"/> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.	
10 <input type="checkbox"/> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 <input type="checkbox"/> Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <input type="checkbox"/> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <input type="checkbox"/> Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.	



OIPE

Does Not Comply
Corrected Diskette Needed

Error on p-3

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002
TIME: 09:37:07

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Output Set: N:\CRF3\05102002\J027736.raw

3 <110> APPLICANT: Presta, Leonard G.
 4 Namenuk, Angela K.
 6 <120> TITLE OF INVENTION: NON-HUMAN PRIMATE Fc RECEPTORS AND METHODS OF USE
 8 <130> FILE REFERENCE: 11669.92US01
 10 <140> CURRENT APPLICATION NUMBER: US 10/027,736
 C--> 11 <141> CURRENT FILING DATE: 2002-04-25
 13 <160> NUMBER OF SEQ ID NOS: 72
 15 <170> SOFTWARE: PatentIn version 3.1
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 20 <213> ORGANISM: Cynomolgus
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 25 <223> OTHER INFORMATION: FcgammaRI alpha-chain
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 33 ttacagtgtg aggtgcccccg tctgcctggg agcagctcca cacagtgggt tctcaatggc 180
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 39 cacagagact ggctactact gcaggtatcc agcagagtct tcacagaagg agaacctctg 360
 41 gccttgaggt gtcatgcattt gaggataag ctgggttaca atgtgcttta ctataaaaat 420
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 45 agtcacaacg ggc当地 atggaaagc atcgctacac atcagcagga 540
 47 gtatctgtca ctgtgaaaga gctatttcca gtc当地 tgc tgaatgc当地 cgtgacatcc 600
 49 cgc当地 ctggggatct ggtc当地 ctggggatct agctgtgaaa caaagttgt tctgc当地 660
 51 cctggtttgc agcttactt ctccctctac atggc当地 agaccctcg aggccaggaaac 720
 53 acgtccctctg aataccaaat actaactgct agaagagaag actctgggtt ttactgggtc 780
 55 gagggccacca cagaagacgg aaatgtccctt aagc当地 ccc ctgagttgga gcttcaagtg 840
 57 cttggccctcc agttaccacat tccctgtctgg cttcatgtcc ttttcttatct ggttagtgggaa 900
 59 ataatgtttt tagtgaacac tggctctgg gtagacaatac gtaaaagaact gaaaagaaaag 960
 61 aaaaagtggaa atttagaat atctttggat tctgctcatg agaagaaggt aacttccagc 1020
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002
TIME: 09:37:07

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 82 ttgcactgtg aggtgcgtca tctgcctggg agcagctcta cacagtggg tctcaatggc 180
 84 acagccactc agacctcgac ccccagctac agaatcacct ctgccagttt caatgacagt 240
 86 ggtgaataca ggtgccagag agtctctca gggcgaagtg accccataca gctggaaatc 300
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 90 gccttgaggt gtcatgcgtg gaaggataag ctgggtaca atgtgcttta ctatcgaaat 420
 92 ggcaaaagcct ttaagttttt ccactggaaat tctaaccctca ccattctgaa aaccaacata 480
 94 agtcacaatg gcacctacca ttgctcaggc atggaaagc atcgctacac atcagcagga 540
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 98 ccactcctgg agggaaatct ggtcaccctgt agctgtgaaa caaagtgcctt cttgcagagg 660
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 104 gaggctgcca cagaggatgg aatgtcctt aagcgcagcc ctgagttgga gcttcagtg 840
 106 cttggcctcc agttaccaac tcctgtctgg tttcatgtcc tttctatct ggcagtggg 900
 108 ataatgttt tagtgaacac tggctctgg gtgacaatac gtaaagaact gaaaagaaaag 960
 110 aaaaagtggg atttagaaat ctctttggat tctggcatg agaagaaggt aatttccagc 1020
 112 cttcaagaag acagacattt agaagaagag ctgaaatgtc aggaacaaaa agaagaacag 1080
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 133 ccgtggatca acgtgtccg ggaggactct gtgactctga cgtgcggggg cgctcacagc 180
 135 cctgacagcg actccactca gtggttccac aatggaaatc gcatccccac ccacacacag 240
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002

TIME: 09:37:07

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Output Set: N:\CRF3\05102002\J027736.raw

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 170 <223> OTHER INFORMATION: FcgammaRIIA
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178 cccccgtgga	tcaacgtgct	ccaggaggac	tctgtgactc	tgacatgcca	gggggctcgc	180
180 agccctgaga	gcgactccat	tcaatgggttc	cacaatggga	atctcatcc	caccacacg	240
182 cagcccagct	acaggttcaa	gccaacaac	aatgacagcg	gggagttacac	gtgccagact	300
184 gcccagacca	gcctcagcga	ccctgtgcat	ctgactgtgc	tttccgaatg	gctgggtctc	360
186 cagacccctc	acctggagtt	ccaggaggga	gaaaccatca	tgctgaggtg	ccacagctgg	420
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190 cgttggatc	ccacccatc	catccacaa	gcaaaaccaca	gtcacagtg	tgattaccac	540
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198 tcagccaatt	ccactgtatcc	tgtgaaggct	gccccattt	agccaccc	acgtcaaatg	780
200 attgccatca	gaaagagaca	acttgaagaa	accaacaatg	actatgaaac	agctgacggc	840
202 ggctacatga	ctctgaaccc	cagggcacct	actgacgatg	ataaaaacat	ctacctgact	900
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223 gctggacac	ctgcagctcc	cccgaaggct	gtgctgaaac	tcgagccccc	gtggatcaac	180
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227 tccactcagt	ggttccacaa	tggaatctc	atccccaccc	acacgcagcc	cagctacagg	300
229 ttcaggcaca	acaacaatga	tagcggggag	tacaggtgcc	agactggcg	gaccagcctc	360
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245 aatcctgacg	aggctgacaa	agttgggct	gagaacacaa	tcacctattc	acttctcatg	840
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251 <211> LENGTH: 876						
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253 <213> ORGANISM: Homo sapiens						

*→ requires elaboration for N
See error summary sheet item 9*

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002

TIME: 09:37:07

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Output Set: N:\CRF3\05102002\J027736.raw

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268	gtgctccagg	aggactctgt	gactctgaca	tgccgggg	ctcacagccc	tgagagcgac	240
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309	gaccgtgtga	ctctgaagt	ccaggggagcc	tactccc	aggacaattc	cacacgg	180
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002

TIME: 09:37:07

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 360 aaagccacac tcaaagacag cggctcctac ttctgcaggg ggcttttgg gagtaaaaat 540
 362 gtgtcttcag agactgtgaa catcaccatc actcaagggtt tggcagtgtc aaccatctca 600
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 366 gtggacacag gactatattt ctctgtgaag acaaaccattc gaagctcaac aagagactgg 720
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 371 <210> SEQ ID NO: 9
 372 <211> LENGTH: 357
 373 <212> TYPE: PRT
 374 <213> ORGANISM: Cynomolgus
 376 <220> FEATURE:
 377 <221> NAME/KEY: MISC_FEATURE
 378 <222> LOCATION: (1)..(357)
 379 <223> OTHER INFORMATION: FcgammaRI 382 <400> SEQUENCE: 9
 384 Met Trp Phe Leu Thr Ala Leu Leu Leu Trp Val Pro Val Asp Gly Gln
 385 1 5 10 15
 388 Val Asp Thr Thr Lys Ala Val Ile Thr Leu Gln Pro Pro Trp Val Ser
 389 20 25 30
 392 Val Phe Gln Glu Glu Thr Val Thr Leu Gln Cys Glu Val Pro Arg Leu
 393 35 40 45
 396 Pro Gly Ser Ser Ser Thr Gln Trp Phe Leu Asn Gly Thr Ala Thr Gln
 397 50 55 60
 400 Thr Ser Thr Pro Ser Tyr Arg Ile Thr Ser Ala Ser Val Lys Asp Ser
 401 65 70 75 80
 404 Gly Glu Tyr Arg Cys Gln Arg Gly Pro Ser Gly Arg Ser Asp Pro Ile
 405 85 90 95
 408 Gln Leu Glu Ile His Arg Asp Trp Leu Leu Leu Gln Val Ser Ser Arg
 409 100 105 110
 412 Val Phe Thr Glu Gly Glu Pro Leu Ala Leu Arg Cys His Ala Trp Lys
 413 115 120 125
 416 Asp Lys Leu Val Tyr Asn Val Leu Tyr Tyr Gln Asn Gly Lys Ala Phe
 417 130 135 140
 420 Lys Phe Phe Tyr Arg Asn Ser Gln Leu Thr Ile Leu Lys Thr Asn Ile
 421 145 150 155 160
 424 Ser His Asn Gly Ala Tyr His Cys Ser Gly Met Gly Lys His Arg Tyr
 425 165 170 175
 428 Thr Ser Ala Gly Val Ser Val Thr Val Lys Glu Leu Phe Pro Ala Pro
 429 180 185 190
 432 Val Leu Asn Ala Ser Val Thr Ser Pro Leu Leu Glu Gly Asn Leu Val

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 05/13/2002
PATENT APPLICATION: US/10/027,736 TIME: 09:37:08

Input Set : A:\Sequence List.txt
Output Set: N:\CRF3\05102002\J027736.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 879

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/027,736

DATE: 05/13/2002

TIME: 09:37:08

Input Set : A:\Sequence List.txt

Output Set: N:\CRF3\05102002\J027736.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:840